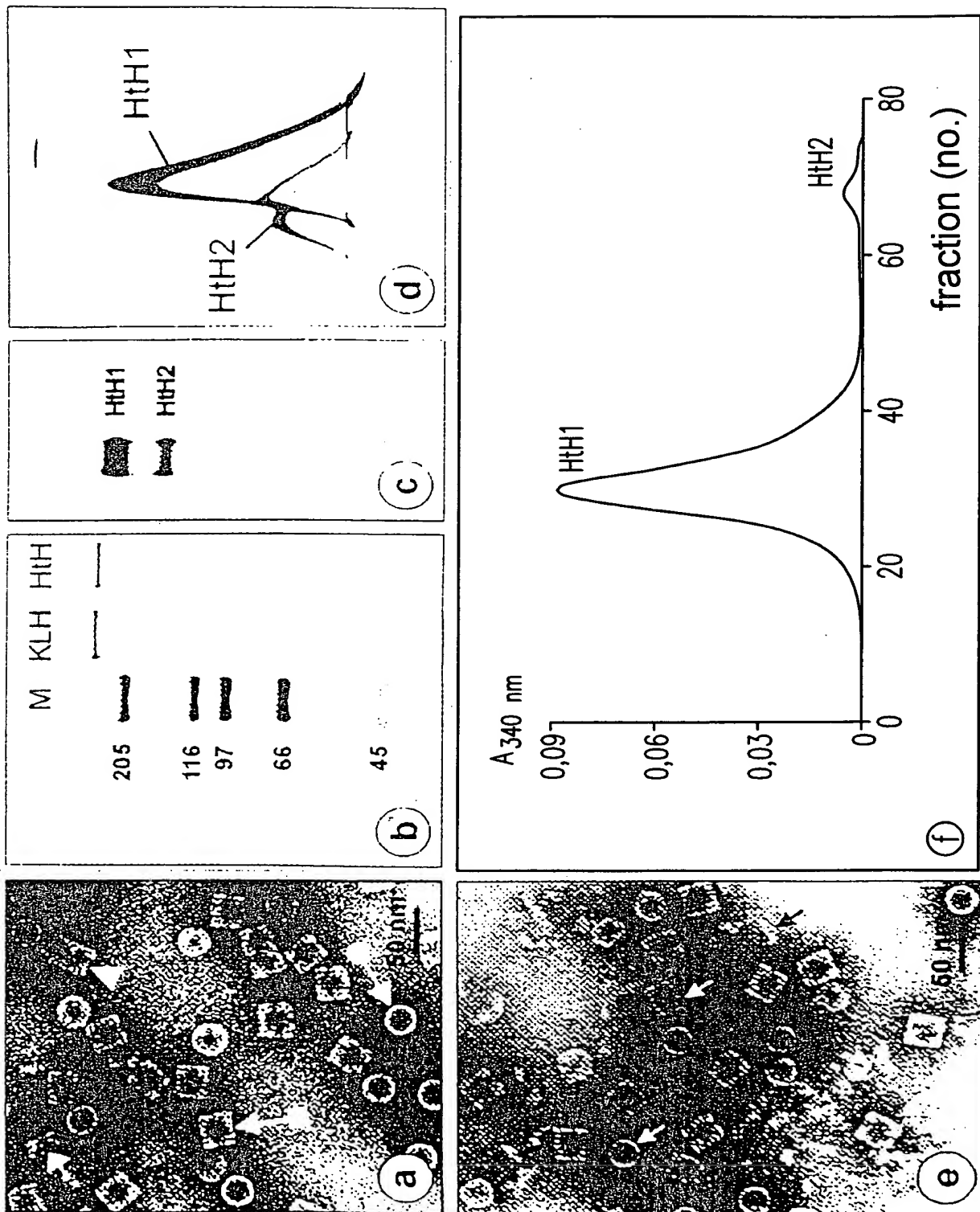
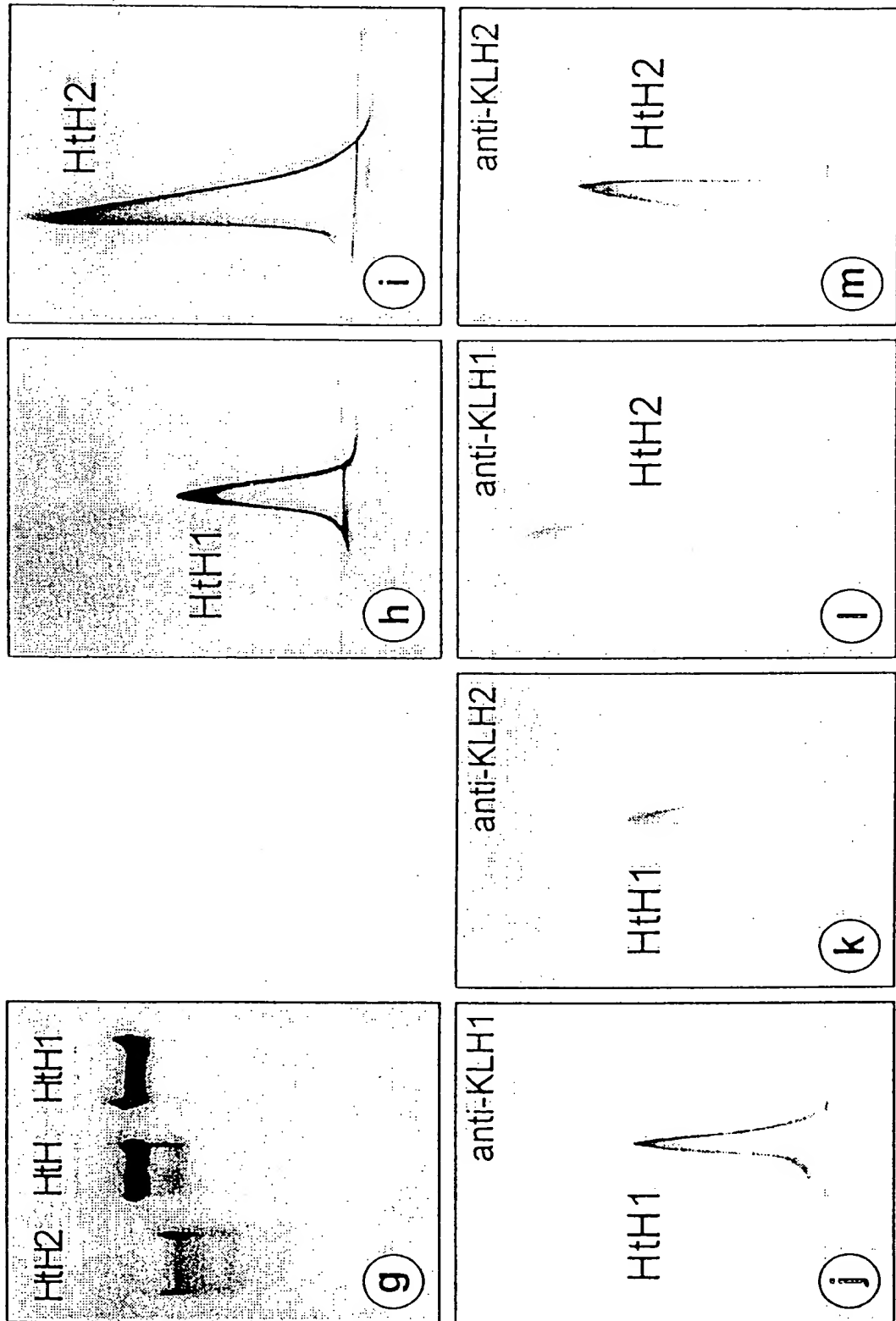


Fig. 1a-f



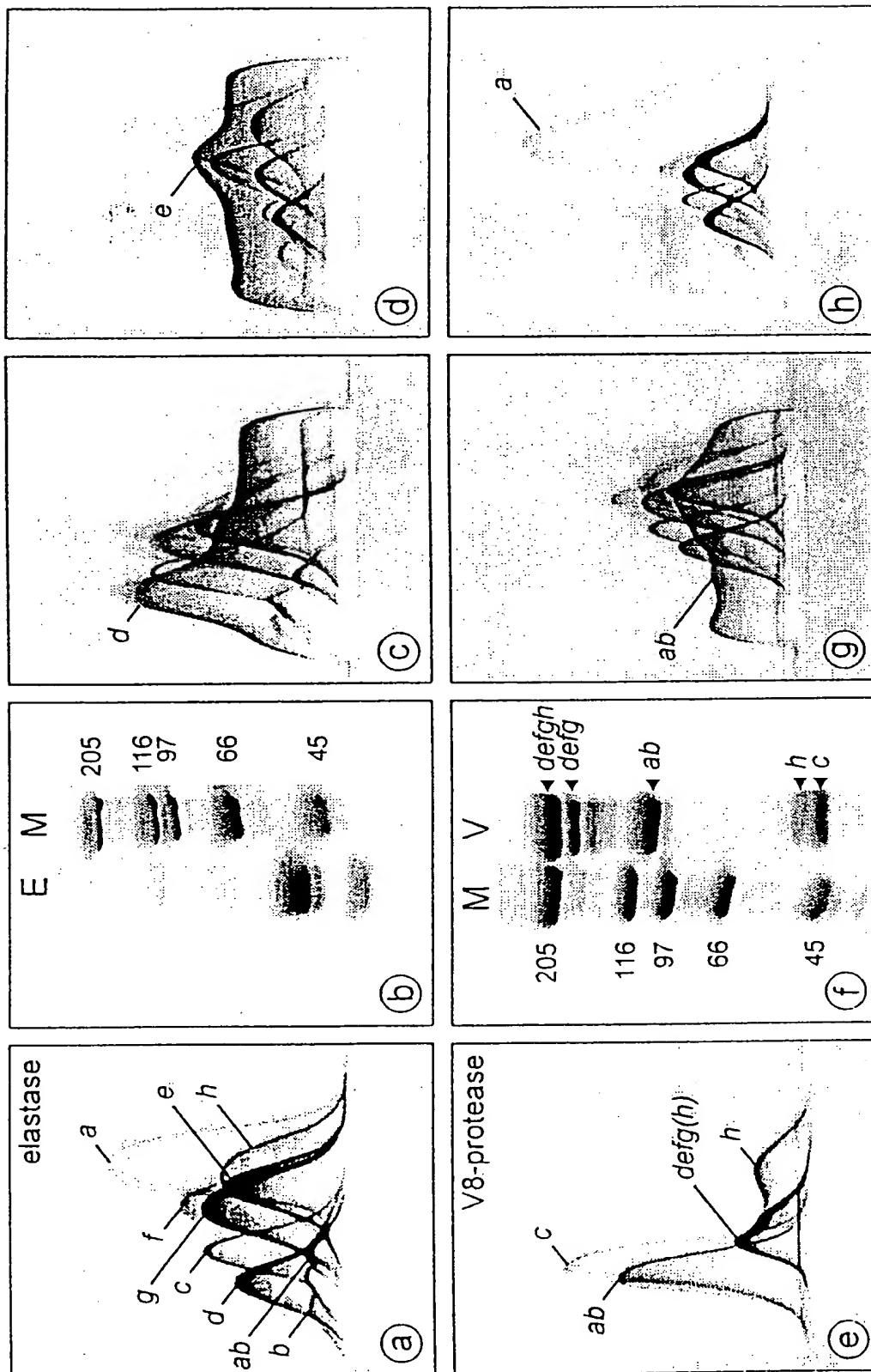
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Fig. 1g-m



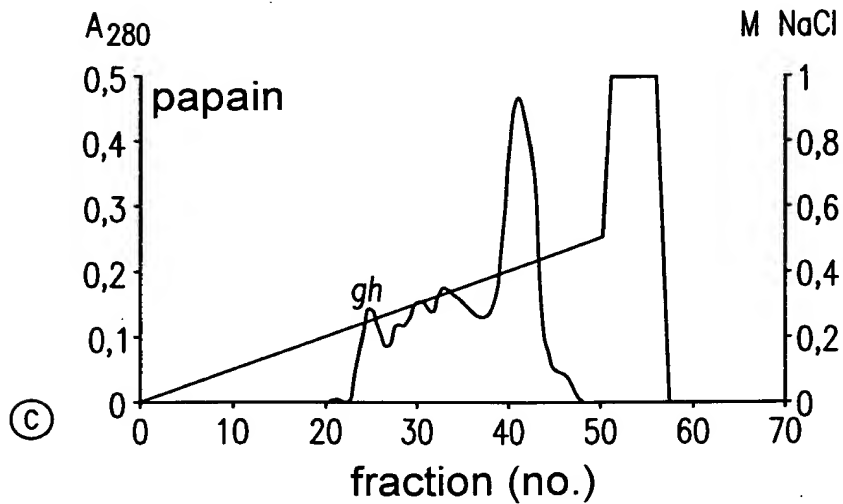
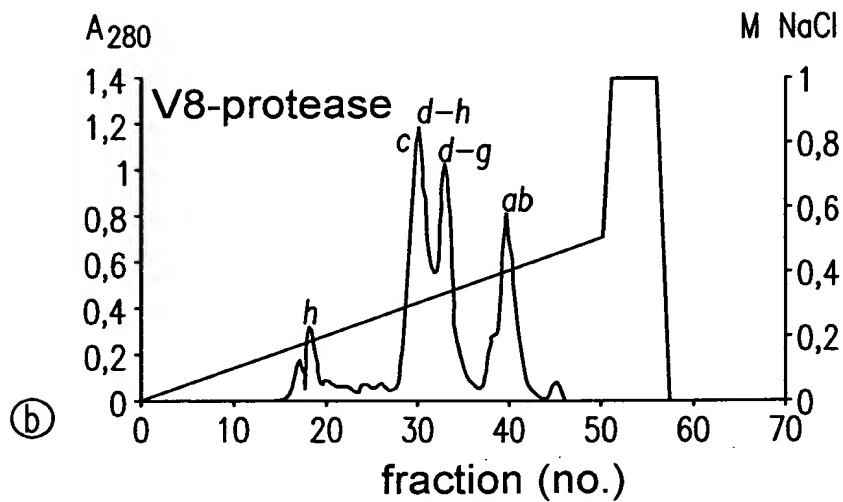
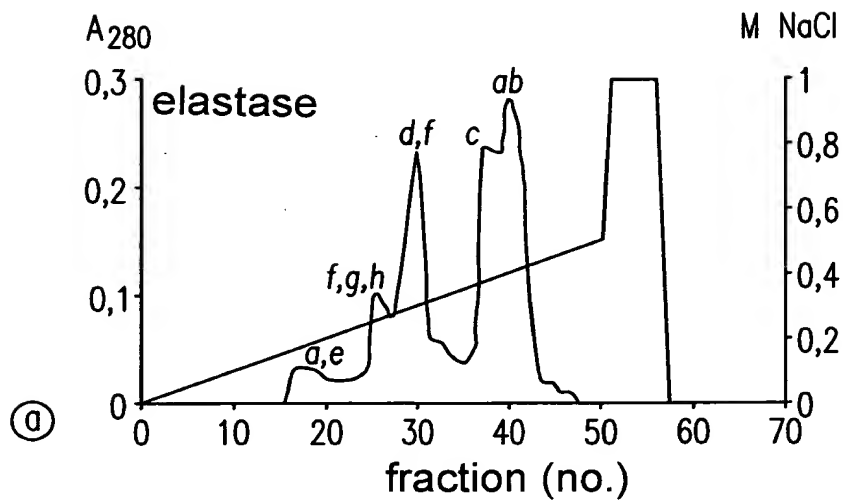
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Fig. 2a-h



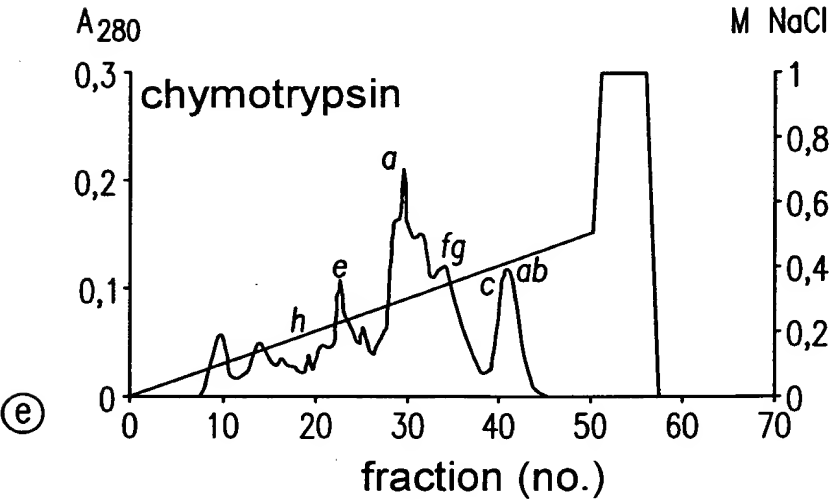
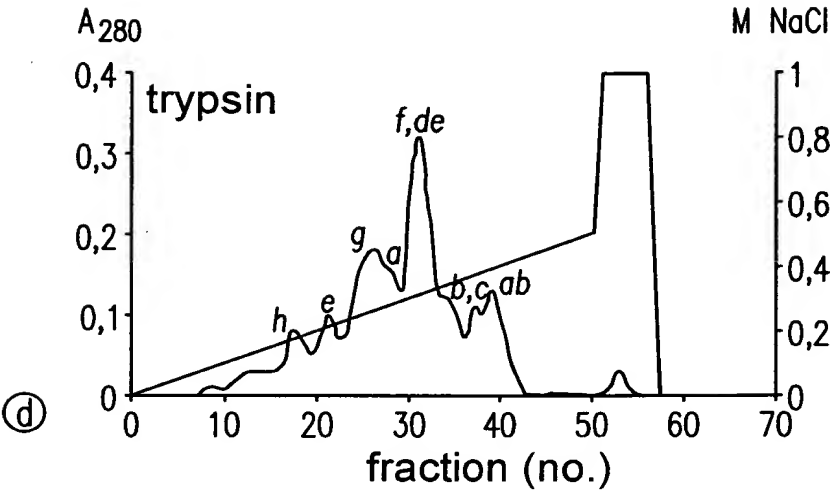
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Fig. 3a-c



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Fig. 3d-e



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Figure 4

HtH1 cDNA sequence and intron structure

Domain a

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Domain b

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Intron b/c

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Domain c

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Intron c/d

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Domain d

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9/29

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Intron d/e

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Domain e

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Intron e/f

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Domain f

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Intron f(1)

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Intron f/g

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REPLACEMENT SHEET NO. 11

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Intron g(2)

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Domain g(2)

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Intron g/h

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Domain h

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3' UTR

TTCACAG

Intron UTR

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3' UTR

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ACACTTTAGAACTTTTAAATGACCTAGAGTGACTTGTAAATATGTAAATATATTCTTCAAAG
ACTCAGCTGAACATATTGTTGGATAACACATCAATTCCTCAACAAATGCTTTATCTTCAC
ATGGATGTATGTAATGTGGCCGGCAATAAAGTATATATATGTATAAAAAAAAAAAAAAAAAA
A

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Figure 5

Derived primary structure of HtH1

Signal peptide

LVQFLLVAGAGA

Domain a

DNVVRKDVSHLTDDDEVQALHGALHDVTASTGPLSFEDITSYHAAPASCDYKGRKIACCVHG
MPSFPFWHRAYVQAEALLSKRKTVMGPYWDWTQTLTHLPSLVTEPIYIDSKGGKAQNTY
WYRGEIAFINKKTARAVDDRLFEKVEPGHYTHLMETVLDALQDEFCKFEIQFELAHNAIH
YLVGGKFEYSMSNLEYTSYDPIFFLHHSNVDRLFAIWQRLQELRGKNPNAMDCAHDLAQ
LQPFNRDSNPVQLTKDHSTPADLFDYKQLGYSYDSLNLNGMTPEQLKTELDERHSKERAF
SFRLSGFGGSANVVYACVPDDPRSDDYCEKAGDFFILGGQSEMPWRFYRPFYDVTEAV
HHLGVPLSGHYVKTFLSVNGTALSPDLLPQPTVAYRPGK

Domain b

GHLDPVHHRHDDDLIVRKNIDHLTREEEYELRMALERFQADTSVDGYQATVEYHGLPARC
PRPDAKVRFACCMHGMASFPHWHRLFVTQVEDALVRRGSPIGVPYWDWTKPMTHLPDLASN
ETYVDPYGHTHHNPFFNANISFEEGHHHTSRMIDSKLFAPVAFGEHSHLFDGILYAFEQED
FCDFEIQFELVHNSIHAWIGGSEDYSMATLHYTAFDPIFYLHHSNVDRLWAIWQALQIRRH
KPYQAHCAQSVEQLPMKPFAPPSPLNNNEKTHSHSVPTDIYDYEEVLHYSYDDLTFGGMNL
EEIEEAHLRQQHERVFAGFLLAGIGTSALVDIFINKPGNQPLKAGDIAILGGAKEMPWAF
DRLYKVEITDSLKTLSDVDGDYEVTFKIHDHMGNALDLDLIPHAADVSEPAH

Domain c

PTFEDEKHSRLRKNVDSLTPETNELRKALELLENDHTAGGFNQLGAFHGEPKWCPNPEA
EHKVACCVHGMVFPWHRLALQAENALRKHGYSYGALPYWDWTRPLSQLPDLVSHEQYTD
PSDHHVKHNPWFNGHIDTVNQDTTRSVREDLYQQPEFGHFTDIAQQVLLALEQDDFCSEV
QYEISHNFIHALVGGTDAYGMASLRYTAYDPIFFLHHSNTDRIWAIWQSLQYRGKPYNTA
NCAIESMRRLQPFGLSSAINPDRTREHAI PFDVFNRYRDNLHYVYDTLEFNGLSISQLDR
ELEKIKSHERVFAGFLLSGIKKSALVKFEVCTPPDNCHKAGEFYLLGDENEMAWAYDRLFK
YDITQVLEANHLHFYDHLFIRYEVFDLKGVS LGTDLFHTANVVHDSGT

Domain d

GTRDRDNYVEEVTGASHIRKNLNDLNTGEMESLRAAFLHIQDDGTYESIAQYHGKPGKCQL
NDHNIACCVHGMPTFPQWHRLYVQVENALLNRGSGVAVPYWEWTAPIDHLPHFIDDATYF
NSRQQRYDPNPFPRGKVTFENAVTTRDPQAGLFNSDYMYENVLLALEQENYCDFEIQFELV
HNAHSMGLGGKGQYSMSLDYSAFDPVFFLHHANTDRLWAIWQELQRFRELPYEEANCAN
LMHQPLKPFSDPHENHDNVTLYSKYPQDGFQYQNHFGYKYNLEFHHLSIPSLDATLKQRR
NHDRVFAFGLLHNIGTSADITIYICLPDGRRGNDCSHEAGTFYILGGETEMPFIFDRLYKF
EITKPLQQLGVKLHGGVFELELEIKAYNGSYLDPHTFDPTIIFEPGT

Domain e

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DTHILDHDHEEEIILVRKNIIDLSPRERVSLVKALQRMKNDRSADGYQAIASFHALPPLCPN
PSAAHRYACCVHGMATFPQWHRLYTVQVQDALRRHGSLVGI PYWDWTKPVNELPELLSSAT
FYHPIRNINISNPFLGADIEFEGPGVHINTERLRFHSGDHDGYHNWFFETVLFALQE
DYCDFEIQFEIAHNGIHTWIGGSVYGMGHLHYASYDPIFYIHHSQTDRIWAIWQELQKYR
GLSGSEANCAIEHMRTPPKPFSFGPPYNLNSHTQEYSKPEDTFDYKKFGYRYDSLELEGRS
ISRIDELIQQRQEKDRTFAGFLLKGFGTSASVSLQVCRVDHTCKDAGYFTILGGS AEMPWA
FDRLYKYDITKTLHDMNLRHEDTFSIDVTITSYNGTVLSGDLIQTPSIIFVPGR

Domain f

HKLNSRKHTPNRVRHELSSLSSRDIA SLKAALTSLOHDNGTDGYQAI AAFHGVPAQCHEPS
GREIACCIHGMATFPHWHRLYT LQLEQALRRHGSSVAVPYWDWTKPITELPHILTDGEYYD
VWQNAVLANPFARGYVKIKDAFTVRNVQESL FKMSSFGKHSLLFDQALLALEQTDYCDFEV
QFEVMHNTIHYLVGGRQTYAFSSLEYSSYDPIFFIHH SFVDKIWAVWQELQSRRLQFRTA
DCAVGLMGQAMRPFNKDFNHSFTKKHAVPNTVFDYEDLGYN YDNLEISGLNLNEIEALIA
KRKSHARVFAGFLLFGLGTSADIHLEICKTSENCHDAGVIFILGGS AEMHWAYNRLYKYDI
TEALQEFDINPEDVFHADEPFFLRLSVVAVNGTVIPSSHLHQPTIIYEPGE

Domain g

DHHDDHQSGS IAGSGVRKDVNTLTKAETDNLREALWGV MADHGPNGFQAIAAFHGKPALCP
MPDGHNYSCCTHGMATFPHWHRLYTKQMEDAMRAHGSHVGLPYWDWTA AFTHLPTLVTDTD
NNPFQHGHI DYLVNSTTRS PRDMLFNDPEHGSESFFYRQVLLALEQTD FCKFEVQFEITHN
AIHSWTGGHSPYGMSTLDFTAYDPLFWLHHSNTDRIWAVWQALQEYRGLPYNHANCEIQAM
KTPLRPFSDDINHN PVTKANAKPLDVFEYNRLSFQYDNLI FHGYSIPELDRVLEERKEEDR
IFAAFLLSGIKRSADVVDICQPEHECVFAGTFAILGGELEMPWSFDRLFRYDITKVMKQL
HLRHDSDFTRVKIVGTDDHELPSDSVKAPTIEFEPG

Domain h

VHRGGNHEDEHDDRLADVLRKEVDFLSLQEANA IKDALYKLQND DSKGGFEA IAGYHGY
PNMCPERGT DKYPCCVHGMPVFPHWHRLHTIQMERALKNHGSPMGI PYWDWTKKMSSLPSF
FGDSSNNNPFYKYYIRGVQHETTRDVNQRLFNQTKFGEFDYLYLT LQVLEENSYCDFEVQ
YEILHNAVH SWLGGTGQYSMSTLEYSAFDPVFMIHHS SLDRIWILWQKLQKIRMKPYALD
CAGDRLMKDPLHPFN YETVNEDEFTRINSFPSILFDHYRFNYEYDNMRIRGQDIHELEEV I
QELRNKDRI FAGFVLSGLRISATVKVFIH SKNDTSHEEYAGEFAVLGGEKEMPWAYERMLK
LDISDAVHKLHVKDEDIRFRVVVTAYNGDVVTTRLSQPFIVHRPAHVAHDILVIPVGAGHD
LPPKVVVKSGTKVEFTPIDSSVNKAMVELGSYTAMAKCIVPPFSYHGFELDKVYSVDHGDY
YIAAGTHALCEQNLRLHIHVEHE

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Figure 6

HtH2 cDNA sequence and intron structure

Domain b

CACAGACTGTTTCGTCACCCAGGTGGAAGATGCTCTGATCAGGCGAGGATCGCCTATAGGGG
TCCCCTACTGGGACTGGACTCAGCCTATGGCGCATCTCCCAGGACTTGCAGACAACGCCAC
CTATAGAGATCCCATCAGCGGGGACAGCAGACACAACCCCTTCCACGATGTTGAAGTTGCC
TTTGAAAATGGACGTACAGAACGTCACCCAGATAGTAGATTGTTTGAACAACCTTTATTTG
GCAACATACGCGTCTCTTCGACAGTATAGTCTATGCTTTTGAGCAGGAGGACTTCTGCGA
TTTTGAAGTTCAATTTGAGATGACCCATAATAATATTACGCCTGGATTGGTGGCGGCGAG
AAGTATTCCATGTCTTCTCTACACTACACAGCCTTCGACCCTATCTTCTACCTTCGTCACT
CCAACACTGACCGGCTCTGGGCAATTTGGCAAGCGTTGCAGATACGAAGAAACAGGCCTTA
CAAGGCTCATTGTGCTTGGTCTGAGGAACGCCAGCCTCTCAAACCTTTGCCTTCAGTTCC
CCACTGAACAACAACGAAAAACCTACGAAAACCTCGGTGCCCACCAACGTTTACGACTACG
AAGGAGTCCTTGGCTATACTTATGATGACCTCAACTTCGGGGGCATGGACCTGGGTCAGCT
TGAGGAATACATCCAGAGGCAGAGACAGAGAGACAGGACCTTTGCTGGTTTCTTTCTGTCA
CATATTGGTACATCAGCGAATGTTGAAATCATTATAGACCATGGGACTCTTCATACCTCCG
TGGGCACGTTTGCTGTTCTTGGCGGAGAGAAGGAGATGAAATGGGGATTGACCGTTTGTA
CAAATATGAGATTACAGATGAACTGAGGCAACTTAATCTCCGTGCTGATGATGTTTTTCAGC
ATCTCTGTTAAAGTAACTGATGTTGATGGCAGTGAGCTGTCCTCTGAACTCATCCCATCTG
CTGCTATCATCTTCGAACGAAGCCATA

Intron b/c

GTAAGTAGCTACCTGTTTATTCAATTTTTTCGCTTTGCCAATCAATTCATTCAGCTTGAAA
TTCAATAATTGTGTTTTGTCATGGCTGAAAACCAATTTGAACTCTTTTCTTTTCTCAGGTGCG
AACTCAAATAAATAATCACTAATTGTTATGCACGCGGGTAGGGCATACTACTATATCCAC
ATCGGTCATCTCAAATGCAAACAAATTGTCTTATTTCCGTTGGGACAAGCAAACCCCTT
TCCTGTAATCTTGCCCTTTGGCATCCACTGGAATTAATGTTGACTGGTAATTGATACTGGCT
CTCTTCTTGTCATAGAGTTAATATCTATAGTTTGTAAATCTTTATGATTTTGCTATTTATAT
TTCGACAGCATGCTATAGACACCCTAGACTATTGTATAGCCACTTGTATTGTTTTTCCATT
TATTATTTATAACAGAACATGGCTTGTAATTTTTATTTACCTTCCAG

Domain c

TTGACCATCAGGACCCGCATCATGACACAATCATTAGGAAAAATGTTGATAATCTTACACC
CGAGGAAATTAATTCTCTGAGGCGGGCAATGGCAGACCTTCAATCAGACAAAACCGCCGGT
GGATTCCAGCAAATTGCTGCTTTTTCACGGGGAACCCAAATGGTGCCCAAGTCCCGATGCTG
AGAAGAAGTTCTCCTGCTGTGTCCATGGAATGGCTGTCTTCCCTCACTGGCACAGACTCCT
GACCGTGCAAGGCGAGAATGCCCTGAGAAAGCATGGATGTCTCGGAGCTCTCCCCTACTGG
GACTGGACTCGGCCCTGTCTCACCTACCTGATTTGGTTTTTGGTAAGTAGCAGAACTACAC
CGATGCCATATTCCACCGTGGAAGCCCGAAACCCCTGGTACAGCGGCCATATTGATACAGT
TGGTGTGACACAACAAGAAGCGTCCGTCAAGAACTGTATGAAGCTCCTGGATTTGGCCAT
TATACTGGGGTCGCTAAGCAAGTGCTTCTGGCTTTGGAGCAGGATGACTTCTGTGATTTTG
AAGTCCAGTTTGAGATAGCTCACAATTTCAATTCACGCTCTTGTCGGCGGAAGCGAGCCATA
TGGTATGGCGTCACTCCGTTACACTACTTATGATCCAATTTTCTACCTCCATCATTCTAAC
ACTGACAGACTCTGGGCTATATGGCAGGCTCTACAAAAGTACAGGGGGCAAACCTTACAATT
CCGCCAACTGCGCCATTGCTTCTATGAGAAAACCCCTACAACCCTTTGGTCTGACTGATGA
GATCAACCCGGATGATGAGACAAGACAGCATGCTGTTCTTTTCAAGTGTCTTTGATTACAAG
AACAACCTTCAATTATGAATATGACACCCTTGACTTCAACGGACTATCAATCTCCCAGCTGG
ACCGTGAAGTGTACGGAGAAAGTCTCATGACAGAGTATTTGCCGGATTTTTGCTGCATGG

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TATTCAGCAGTCTGCACTAGTTAAATTCTTTGTCTGCAAATCAGATGATGACTGTGACCAC
TATGCTGGTGAATTCTACATCCTTGGTGATGAAGCTGAAATGCCATGGGGCTATGATCGTC
TTTACAAATATGAGATCACTGAGCAGCTCAATGCCCTGGATCTACACATCGGAGATAGATT
CTTCATCAGATACGAAGCGTTTGATCTTCATGGTACAAGTCTTGGAAGCAACATCTTCCCC
AAACCTTCTGTCTACATGACGAAGGGGCAG

Intron c/d

GTGAGAACATTGATAATAGTTCAAATgAAGTATATCCGATTCAAGCTGTCGATACAAGATg
AGATACATAATCACAATGTTTGTATTAGATATCTCTCTTAATTTAATGCCGCTTTTATCAA
TATTCGAGCAATCCTTCAGCAACATACACCAGCAAATGTTTCATCAACAGACTATATTATT
TAATCTTTTAAAAATCCTTTTCTGTTGTTATAAATACTTAAAGTATCGAATTCCTTGAATG
CGTCTTCTCTGCAGCATATAGTTAAGTTGTTGTGTTTCTCTGTCTGTCAG

Domain d

GTCACCATCAGGCTGACGAGTACGACGAAGTTGTAAGTCTGCAAGCCACATCAGAAAGAA
TTTAAAAGATCTGTCAAAGGGAGAAGTAGAGAGCCTAAGGTCTGCCTTCCTGCAACTTCAG
AACGACGGAGTCTATGAGAATATTGCCAAGTTCCACGGCAAGCCTGGGTTGTGTGATGATA
ACGGTCGCAAGGTTGCCTGTTGTGTCCATGGAATGCCACCTTCCCCCAGTGGCACAGGCT
CTATGTCCTCCAGGTGGAGAATGCTTTGCTGGAGAGAGGATCTGCCGTCTCTGTGCCATAC
TGGGACTGGACTGAAACATTTACAGAGCTGCCATCTTTGATTGCTGAGGCTACCTATTTCA
ATTCCCGTCAACAAACGTTTGACCCTAATCCTTTCTTCAGAGGTAAAATCAGTTTTGAGAA
TGCTGTTACAACACGTGATCCCCAGCCTGAGCTGTACGTTAACAGGTACTACTACCAAAC
GTCATGTTGGTTTTTTGAACAGGACAACACTACTGCGACTTCGAGATACAGTTTTGAGATGGTTC
ACAATGTTCTCCATGCTTGGCTTGGTGGAAGAGCTACTTATTCTATTTCTTCTCTTGATTA
TTCTGCATTTCGACCCTGTGTTTTTTCCTTCACCATGCGAACACAGATAGATTGTGGGCCATC
TGGCAGGAGCTGCAGAGGTACAGGAAGAAGCCATACAATGAAGCGGATTGTGCCATTAACC
TAATGCGCAAACCTCTACATCCCTTCGACAACAGTGATCTCAATCATGATCCTGTAACCTT
TAAATACTCAAAACCCACTGATGGCTTTGACTACCAGAACAACCTTTGGATACAAGTATGAC
AACCTTGAGTTCAATCATTTTCAGTATTTCCAGGCTTGAAgAAATCATTTCGtATTAGACAAC
GTCAAGATCGTGTGTTTGCAGGATTCTCTCCTTCACAACATTGGGACATCCGCAACTGTTGA
GATATTCTGTCTGTGTCCCTACCAACGCGGTGAGCAAACCTGTGAAAACAAAGCCGGAACA
TTTGCCGTACTCGGAGGAGAAACAGAGATGGCGTTTCATTTTGACAGACTCTACAGGTTTG
ACATCAGTGAAACACTGAGGGACCTCGGCATACAGCTGGACAGCCATGACTTTGACCTCAG
CATCAAGATTCAAGGAGTAAATGGATCCTACCTTGATCCACACATCCTGCCAGAGCCATCC
TTGATTTTTGTGCCTGGTTCAAGT

Intron d/e

AAGAAAGTTTCACTGTCTAAATCTTTTTTTTATGATAGAGGGTAGAGAAGTGGAGACAATGT
GACAATATATTGAATAAAGTTGTTTAAAATTTATAACTCTCATAAGTTTCATATTATGCTGA
AGCTGTAGCCATCTATAACTGTGTAACATGAAATGTTAAGACATTAACCTAAATACTTCAG
CTGATAACAAAACAATGTTAATACATACGTCAATGTAACATTTTCTTATCTTTAGGTTATA
GCATAAACACTTCAGAGATACAGTGACGAAAACCTCTATTTAAATATTTTCAGGT

Domain e

TCTTTCCTGCCTCCTGATGGGCATTTCAGATGACATCCTTGTGAGAAAAGAAGTGAACAGCC
TGACAACCAAGGGAGACTGCATCTCTGATCCATGCTCTGAAAAGTATGCAGGAAGACCATTC
ACCTGACGGGTTCCAAGCCATTGCCTCTTTCCATGCTCTGCCACCACTCTGCCCTTCACCA
TCTGCAGCTCACCGTTATGCTTGCTGTGTCCACGGCATGGCTACATTTCCCCAGTGGCACA
GATTGTACACTGTACAGTTCCAGGATGCACTGAGGAGACATGGAGCTACGGTAGGTGTACC
GTATTGGGATTGGCTGCGACCGCAGTCTCACCTACCAGAGCTTGTCAACCATGGAGACATAC

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CATGATATTTGGAGTAACAGAGATTTCCCAATCCTTTCTACCAAGCCAATATTGAGTTTG
AAGGAGAAAACATTACAACAGAGAGAGAAGTCATTGCAGACAACTTTTTGTCAAAGGTGG
ACACGTTTTTTGATAAACTGGTTCTTCAAACAAGCCATCCTAGCGCTGAGCAGGAAAACACTAC
TGTGACTTTGAGATTGAGTTTGAATTTCTTCAACACGGCGTTCACACGTGGGTTCGGAGGCA
GTCGTACCTACTCTATCGGACATCTTCATTACGCATTCTACGACCCTCTTTTCTACCTTCA
CCATTTCCAGACAGACCGTATTTGgGCAATCTGGCAAGAACTCCAGGAACAGAGAGGGCTC
TCGGGTGATGAGGCTCACTGTGCTCTCGAGCAAATGAGAGAACCATTGAAGCCTTTTCAGCT
TCGGCGCTCCTTATAACTGGAATCAGCTCACACAGGATTTCTCCCGACCCGAGGACACCTT
CGACTACAGGAAGTTTGGTTATGAATATGACAATTTAGAATTCCTGGGAATGTCAGTTGCT
GAACTGGATCAATACATTATTGAACATCAAGAAAATGATAGAGTATTCGCTGGGTTCCTGT
TGAGTGGATTTCGGAGGTTCCGCATCAGTTAATTTCCAGGTTTGTAGAGCTGATTCCACATG
TCAGGATGCTGGGTACTTCACCGTTCTTGGTGGCAGTGCTGAGATGGCGTGGGCATTTGAC
AGGCTTTTACAAATATGACATTACTGAACTCTGGAGAAAATGCACCTTCGATATGATGATG
ACTTCACAATCTCTGTGAGTCTGACCGCCAACAACGGAAGTGTCTTGAGCAGCAGTCTAAT
CCCAACACCGAGTGTTCATATTCAGCGGGGACATC

Intron e/f

AAGTAGTAAACTGCTCAGATTGTTTTTCATAATTACTCCACTATTAAGTAAAAAGTACTAGT
AATTCAATAGTACTGTTTACAGAGAAATGTAACACAATAGACCACAGAGTCCATTTGTAA
ACGCCTTTGGCTTGTAAGTCTGAGGTTTTGGTGACTGATGGAAAGCTAAAATATATTTTG
ACAG

Domain f(1)

GTGACATAAATACCAGGAGCATGTCACCGAACCGTGTTGCGCGTGAGCTGAGCGATCTGTC
TGCGAGGGGACCTGTCTAGTCTCAAGTCTGCTCTGCGAGACCTACAGGAGGATGATGGCCCC
AACGGATACCAGGCTCTTGACGCCTTCCATGGGCTACCAGCAGGCTGCCATGATAGCCGGG
GAAATGAGAT

Intron f

ATATTTAAAGTATTTTATCTTACGCATGACCCTGACCCTATTATTTTTTTAATCCTATGAT
GAAACATTTACTTAGACTGGCTTGTGAGCCCCAGGCAAAATGCACTGTAAAAATACACTGA
CAGAGGATTAGGCATTCTTGGGAGTACTGTATAGTTAGTTGCATACATATTAGCGTTCCT
CACTAAAACGAATCTCTGAATGCTATCAATTAAAGATCATGATGCTTTGATTGTGTCTACT
GTATTTAAATGGTGTAAAGATTTGCAATTACAATATACACAAACACGTTTCCTGCATCTC
GGAGAATGCAATCTTTCGTTGTACGCGTCTGTTTTTCATATTTTTATGCATGTAGTTTGCAC
TACTTAGCGTCCAATAAATCCATTACAAAATCACACAAACAAACGATTTTAGGAATGTGA
CTGTAGCTGCAACGAATATACCTGATCCTTTCTTGTTCCAGAT

Domain f(2)

CGCATGTTGCATTACAGGGATGCCGACCTTCCCCAGTGGCACAGACTGTACACCCTGCAG
TTGGAGATGGCTCTGAGGAGACATGGATCATCTGTGCGCATCCCCTACTGGGACTGGACAA
AGCCTATCTCCGAACCTCCCTCGCTCTTACCAGCCCTGAGTATTATGACCCATGGCATGA
TGCTGTGGTAAACAACCCATTCTCCAAAGGTTTTGTCAAATTTGCAAATACCTACACAGTA
AGAGACCCACAGGAGATGCTGTTCCAGCTTTGTGAACATGGAGAGTCAATCCTCTATGAGC
AACTCTTCTTGCTCTTGAGCAAACCGACTACTGTGATTTTGAAGTACAGTTTGAAGTCCT
CCATAACGTGATCCACTACCTTGTGGTGGACGTCAGACCTACGCATTGTCTTCTCTGCAT
TATGCCTCCTACGACCCATTCTTCTTTATACACCATTCTTTTCTGGATAAGATGTGGGTAG
TATGGCAAGCTCTTCAAAGAGGAGGAACTTCCATACAAGCGAGCTGACTGTgcTGTCAA
CCTAATGACTAAACCAATGAGGCCATTTGACTCCGATATGAATCAGAACCCATTACAAAG
ATGCACGCAGTTCCCAACACACTCTATGACTACGAGACACTGTACTACAGCTACGATAATC
TCGAAATAGGTGGCAGGAATCTCGACCAGCTTCAGGCTGAAATTGACAGAAGCAGAAGCCA

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CGATCGCGTTTTTCTGGATTCTTGCTTCGTGGAATCGGAACTTCTGCTGATGTCAGGTTT
TGGATTTGTAGAAATGAAAATGACTGCCACAGGGGTGGAATAATTTTCATCTTAGGTGGAG
CCAAGGAAATGCCATGGTCATTTGACAGAACTTCAAGTTTGATATCACCCATGTACTCGA
GAATGCTGGCATTAGCCCAGAGGACGTGTTTGATGCTGAGGAGCCATTTTATATCAAGGTT
GAGATCCATGCTGTAAACAAGACCATGATACCGTCGTCTGTGATCCCAGCCCCAACTATCA
TCTATTCTCCTGGGGGAAAG

Intron f/g

GTGAGAGAACCAGTAATAGCTACTGTCTACAAAGAATGTGTTTCATTTAAAGACCTGACTGT
AGGCCGATGGCTGCTGTCATCTCCTCCGCCCTCCTCCTCCTGTTCCCTCCGAAGGGGTCA
GTTTCAGGTTCTCTTGCCAATATGCCAAGCAGACCTCCTGAGCAGGCAGTATATATACGTA
AGGGAAGCAAGTATGGACCATCGCGCGGCATGTAGAGATACAATGATCAGCTGTCTGTCTGT
TCCACTCCTGTCAGACAATGAGATAAACATGAATACAGTATTACTCAGCAGCGTTCCAATT
TTCAACCCTCGTATTTATTAAAAAAGGAATTTTAAATATATTTTCTCCTTGTGAAATA
TTTTAGTAACGTGTTAATCGATATAGAGTGGAGTAGTGACGCTTTATTTTCGGTTCATTCTCG
AAACAAAAATAATAAGTCCACTGAACTCTCTTAAATTGTTTTTACAACCTTCAACTGCCA
CAGACGTAATCCCTCACGTTATTTTGAGCTGACAACGTGTTGAATTGAGTGTGTTCCGAAT
TCTAAATAAGCATGTATATATTTACGTCTCATGCAAGTAATATATGTTTAACTGATGACGT
CACTTGGTGACCACTGATTTAGTTTCCTTTGTCTAATTGCAGTTTCTGTTGTCACGGGGAC
GGTGGGGAAGCCAGGTTCTCCTGTCACGCTGAATATCCCGTTCGAATCCCCCACATGGGT
ACAAAGTGTGATGCCTATTTCTGGTGTCCCCCACCCTGATATTGCTGGAATAAGTGGCTTA
ATACCATATACACTCACTCTATTGTACACTACTGCCACCGGCTCACACCTCTGATGCTTC
TGTTCTATCCAG

Domain g(1)

GTGCGCTGCTGACAGTGCGCACTCTGCCAACATTGCTGGCTCTGGGGTGAGGAAGGACGT
CACGACCCTCACTGTGTCTGAGACCGAGAACCCTAAGACAGGCTCTTCAAGGTGTCATCGAT
GATACTGGTCCCAATGGTTACCAAGCAATAGCATCCTTCCACGGAAGTCCTCCAATGTGCG
AGATGAACGGCCGCAAGGTTGCCTGTTGTGCTCAG

Intron g(1)

GTAATTAATGGATGTGAAGTCAATGTCCGAGGGTATAATAAGGATTTAAATACTTCAGTCG
TGTAATACTGTATGACATGTGTATTGGATGGTGTAGGTATTACAGGTTATAAGGCCAGTGT
GTGTTGGGACGGTTACTTTCTGCACTAGTAATAAGCATTGTATTTAGCTAGCTTTTATCA
TATAACTTTAGTTTCAGGTTTGtGGCAATTGAAATCGAAATTTTCTTTTCAATTTCAAGGTTA
TCGCACTCGTGTGTNAGAATAGTTACTATGCTGCATTGAGAATAACACTATAGTAATAAAG
CATATCATAACAGTAAGAATAACACTATAGTAATAAAGTATATCATNCAGTAAGAATGTCAT
TGTATGATAAATAGGTTATCACACTCGTGTGTTTTAGAATGGTTACTATCCCAGGAATAAC
CACTATGTATTACATGTATATTGGGCAGTGTAAGTAGTAGCATTGTATATTAAATCAGTAT
ATCGTGCTTCAAAACACCAGGATATATGGGGTATACAGTGGGCAGTGTAAGTAGCAACATT
GTATATTAAATCAGTATATCGTACTTCAAAACACCAGGATTATGGGGTATACAGTGGGCAG
TGTAAGTAGTAGCATTGTATATTAAATCAGTATATCGTACTTCAAAACACCAGGATATAAT
TCAGTATATCGTGCTTCAAAACACCAGGATATAATTCAGTATATCGTGCTTCAAAACACCA
GGATATATGGGATATACAGTGCGGGTTTGCATACAACCTCCACCTTTACAG

Domain g(2)

GTATGGcCTCCTTCCcACACTGGCACAGACTGTATGTGAAGCAGATGGAAGATGCCCTGGC
TGACCACGGGTACATATCGGCATCCCTTACTGGGACTGGACAACCTGCCTTCACAGAGTTA
CCCGCCCTTGTCACAGACTCCGAGAACAATCCCTTCCATGAG

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19/29

Intron g(2)

GTCAGTTTAGTCTCCTGTCTGAGCTAACGATACCAATTTCTATTTTCGAGAACCACGATG
ACGAGAAAACAAGCAATATAGATATAGATGCAGTATAGATCAAGTTAATGAATTCATTGCT
ATATGTTTGCTTGTAATAAACTTTAAGAAAACGAGAGCATGCACACAAATGAAACAAACAA
TTATGTGTTTGATAGGAATATGATATATGTATTTGGGGGCTGACGTGAGCAGGGTTGAAGG
GACAGTTTACATTGTGAGTAACACTGGGAGTATTCTTTGATCCACAATATATAGTTTTCATT
GTGTTTCAGCAGTTACAACATAACATTATATCATACATTACGTCGTAACATGCTTCTTTTGTC
CTCTTTTGCCAG

Domain g(3)

GGTCGCATTGATCATCTCGGTGTAACCACGTCACGTTCCCCCAGAGACATGCTGTTTAAACG
ACCCAGAGCAAGGATCAGAGTCGTTCTTCTATAGACAAGTCCTCCTGGCTTTGGAGCAGAC
TGACTACTGCCAGTTCGAAGTCCAGTTTGAGCTGACCCACAACGCCATTCACTCCTGGACA
GGTGGACGTAGCCCTTACGGAATGTCGACCCCTCGAGTTTACAGCCTACGATCCTCTCTTCT
GGCTTCACCACTCCAACACCGACAGAATCTGGGCTGTCTGGCAAGCACTGCAGAAATACCG
AGGACTCCCATAACAACGAAGCACACTGTGAAATCCAGGTTCTGAAACAGCCCTTGAGGCCA
TTCAACGATGACATCAACCACAATCCAATCACCAAGACTAATGCCAGGCCTATCGATTTCAT
TTGATTATGAGAGGTTTAACTATCAGTATGACACCCTTAGCTTCCATGGTAAGAGCATCCC
TGAAGTGAATGACCTGCTCGAGGAAGAGAAAAAGAGAGAGAACATTTGCTGcCTTCCTT
CTTCGTGGAATCGGTTGCAGTGCTGATGTCGTCTTTGACATCTGCCGgCCCAATGGTGACT
GTGTCTTTGCAGGAACCTTTGCTGTGCTGGGAGGGGAGCTaGAAATGCCTTGGTCCTTCGA
CAGACTGTTCCGCTATGACATCACCAAGAGTCATGAATCAGCTCCATCTCCAGTATGATTCA
GATTTTCAGTTTCAGGGTGAAGCTTGTTGCCACCAATGGCACTGAGCTTTCATCAGACCTtC
TCAAGTCACCAACAATTGAACATGAACTTGG

Intron g/h

GTATGTTATCTTATCATCAAATGTGTGATCAGATACTGGAGACGTTTTTCATATTAACCTGG
TCAGCATTAGTTGATGATTTTGGTGCGATGTTGACGACAAGGAGTCAAGCATTAAACACATT
CAACACATCTTTAATCTGATATGAGAAGGGAATAAATTGATCCAGTATTGATGATTGAAGT
TAGATTAACAGTGAAAGATATACCAGTTTTGATAATCGTATAAAACAGTAGCAGAATTGTA
TCGTGAAACTAAATGTGGGAAGGCGAACGCCAAGCAGATTTTAGATTACGATCGTGCTGT
AGAATAATTACAAATAACCCAGACGTCGGAAATGTGGTTGTCTATGGCAATGGTTACGATT
AATTGCTAACATGCACGATTTACCTATTTTACG

Domain h

AGCCACAGAGGACCAGTTGAAGAAACAGAAGTCACTCGCCAACATACTGACGGCAATGCA
CACTTTTCATCGTAAGGAAGTTGATTGCTGTCCCTGGATGAAGCAAACAACCTTGAAGAATG
CCCTTTACAAGCTACAGAACGACCACAGTCTAACGGGATACGAAGCAATCTCTGGTTACCA
TGGATACCCCAATCTGTGTCCGGAAGAAGGCGATGACAAAATACCCCTGCTGCGTCCCCGG
ATGGGCATCTTTCTTACTGGCACAGACTCTTGACCATTCAACTGGAAAGAGCTCTTGAGC
ACAATGGTGCCTGCTTGGTGTTCTTACTGGGACTGGAACAAGGACCTGTCGTCACTGCC
GGCGTTCTTCTCCGACTCCAGCAACAACAATCCCTACTTCAAGTACCACATCGCCGGTGTT
GGTCACGACACCGTCAGAGAGCCAACTAGTCTTATATATAACCAGCCCCAAATCCATGGTT
ATGATTATCTCTATTACCTAGCATTGACCACGCTTGAAGAAAACAATTACTGGGACTTTGA
GGTTCAGTATGAGATCCTCCACAACGCCGTCCACTCCTGGCTTGGAGGATCCCAGAAGTAT
TCCATGTCTACCCTGGAGTATTCGGCCTTTGACCCTGTCTTTATGATCCTTCACTCGGGTC
TAGACAGACTTTGGATCATCTGGCAAGAACTTCAGAAGATCAGGAGAAAGCCCTACAACCT
CGCTAAATGTGCTTATCATATGATGGAAGAGCCACTGGCGCCCTTCAGCTATCCATCTATC
AACCAGGACGAGTTCACCCGTGCCAACTCCAAGCCTTCTACAGTTTTTGACAGCCATAAGT

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TCGGCTACCATTACGATAACCTGAATGTTAGAGGTCACAGCATCCAAGAACTCAACACAAT
CATCAATGACTTGAGAAACACAGACAGAATCTACGCAGGATTTGTTTTGTCAGGCATCGGT
ACGTCTGCTAGTGTCAAGATCTATCTCCGAACAGATGACAATGACGAAGAAGTTGGAACCT
TCACTGTCCTGGGAGGAGAGAGGGGAAATGCCATGGGCCTACGAGCGAGTTTTCAAGTATGA
CATCACAGAGGTTGCAGATAGACTTAAAATTAAGTTATGGGGACACCCTTTAACTTCCGGA
ACTGGAGATCACATCCTTACGAATGGAATCGGTGGTAAACAAGAGCCTACCCAAATCCTTT
CATCATCTACAGACCTGCCAATCATGACTACGATGTTCTTGTTATCCAGTANGGAAGAAA
CCTTCACATCCCTCCCAAAGTTGTCGTCAAGAAAGGCACCCGCATCGAGTTCCACCCAGTC
GATGATTCAGTTACGAGACCAGTTGTTGATCTTGGAAGCTACACTGCACTCTTCAACTGTG
TGGTACCACCGTTCACATACCACGGATTGCAACTGAACCACGTCTATTCTGTCAAGCCTGG
TGACTIONTATGTTACTGGACCCACGAGAGACCTTTGCCAGAATGCAGATGTCAGGATTCAT
ATCCATGTTGAGGATGAGTAA

3' UTR

CGCAACAGGT

Intron UTR

GAGATAAGAAACCCTTCTAACAGTAATACGACACCACATTACAGCTTAAACATGATTGCCA
TCGATGTTTTTCATGTGTAGTATACGCTTTTCAGTTCTACATAATTTTGTTTTTCAAATCAA
GTTTAGCAAATGAATCTATCACTGGAAAATAGGGTAGGGTAGCCAAGTGGTTAAAGCGGTC
ACTGATCACGCCAAAGACGAGTGTCTAACCTGCATGGGTACAAAAGTGAAGACCATTGCT
GGTGTCTACCGCCGTAATATTGTTTTTAGTATTGCTAAAACTTATACTCACCCATGCGCTG
TAAAAGTGAATAATAATCATATTTCAACAAAAGCACAAAACCATTTTCATTTTCATGAAAG
CCTCTTGTTTCACCTGAAAGACGCAAGAGAACAAATAGTTCCCTAACATTATTTTCAGACATTG
GAAATGTCCTGCACGTGTAAACCATATATCCTTTGAAATTTTACGACTGCATCGTATACA
ATTTATGATATAAATTTAAAACTTTAT

3' UTR

TTCTTGGTCTCCACATATTCACATATCAGCACCAAATGGTTTCGAAGGACATTGGCGTTCT
TCTCTGGCAATGCATTTCAATACAACATTGAAAATGACTTCAGCATATCAGTGTGCTTCGA
ACGTGTTCCGGAAGTACTCAAATGTGCTATGACTGAATTATTGTACATACATAACTTATTG
ATGTTCAATAAATAAATGTTGAAACGAAAAAAAAAAAAAAAAAAAAAAAAA

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Figure 7

Derived primary structure of Hth2

Domain b

HRLFVTQVEDALIRRGSPIGVPYWDWTQPM AHLPLGLADNATYRDPI SGDSRHNPFDHDEVA
FENGRTERHPDSRLFEQPLFGKHTRLFDSIVYAFEQEDFCDFEVQFEMTHNNIHAWIGGGE
KYSMSSLHYTAFDPIFYLRHSNTDRLWAIWQALQIRRNRPYKAHCAWSEERQPLKPFASFSS
PLNNNEKTYENSVPTNVYDYEGLVGYTYDDLNF GGMDLGQLEEIYIQRQRQDRTFAGFFLS
HIGTSANVEIIIDHGTLHTSVGTFAVLGGEKEMKWGFDRLYKYEITDELRLQLNLRADDVFS
ISVKVTDVDGSELSSSELIPSAIIIFERSH

Domain c

IDHQDPHHDITIIRKNVDNLTPEEINSLRRAMADLQSDKTAGGFQQIAAFHGEPKWCPSPDA
EKKFSCCVHGMVFPWHRLTLVQGENALRKHGCLGALPYWDWTRPLSHLPDLVLVSSRTT
PMPYSTVEARNPWYSGHIDTVGVDTTRSVRQELYEAPGFGHYTGVAQVLLALEQDDFCDF
EVQFEIAHNFIHALVGGSEPYGMASLRYTTYDPIFYLRHSNTDRLWAIWQALQKYRGKPYN
SANCAIASMRKPLQPFGLTDEINPDDETRQHAPVFSVFDYKNNFNIEYDTLDFNGLSISQL
DRELSRRKSHDRVFAGFLLHGIQQSALVKFFVCKSDDDCDHYAGEFYILGDEAEMPWGYDR
LYKYEITEQLNALDLHIGDRFFIRYEAFFDLHGTSLSGNI FPKPSVIHDEGA

Domain d

GHHQADEYDEVVTAASHIRKNLKDLSKGEVESLRS AFLQLQNDGVYENIAKFHGKPGLCDD
NGRKVACCVHGMPTFPQWHRLYLVLQVENALLERGS AVSVPYWDWTETFTELPSLIAEATYF
NSRQOTFDPNPFPRGKISFENAVTTRDPQPELYVNRYYYQNVMLVFEQDNYCDFEIQFEMV
HNVLHAWLGGRATYSSISLDYSAFDPVFFLHHANTDRLWAIWQELQRYRKKPYNEADCAIN
LMRKPLHPFDNSDLNHDPTVTFKYSKPTDGFQYQNNFGYKYDNLEFNHFSIPRLEETIRIRO
RQDRVFAGFLLHNI GTSATVEIFVCVPTTSGEQNCENKAGTFAVLGGETEMAFHFDRLYRF
DISETLRDLGIQLDSHDFDLSIKIQGVNGSYLDPHILPEPSLIFVPGSS

Domain e

SFLRPDGHSDDILVRKEVNSLTRETASLIHALKSMQEDHSPDGFQAIASFHALPPLCPSP
SAAHRYACCVHGMATFPQWHRLYTVQFQDALRRHGATVGVPYWDWLRPQSHLPDLVTMETY
HDIWSNRDFPNPFYQANIEFEGENITITEREVIADKLFVKGGHVFDKLVLOTSHPSAEQENY
CDFEIQFEILHNGVHTWVGGSRTYSIGHLHYAFYDPLFYLRHHFQTDRIWAIWQELQEQRGL
SGDEAHCALEQMREPLKPFSGAPYNWNQLTQDFSRPEDTFDYRKFGYEYDNLEFLGMSVA
ELDQYIIHQENDRVFAGFLLSGFGGSASVNFQVCRADSTCQDAGYFTVLGGS AEMAWAFD
RLYKYDITETLEKMHLRYDDDFITISVSLTANNGTVLSSSLIPTPSVIFQRGH

Domain f

RDINTRSMSPNVRRELSDLSARDLSSLKSALRDLQEDDGPNGYQALAAFHGLPAGCHDSR
GNEIACCIHGMPPTFPQWHRLYTLQLEMALRRHGSSVAIPYWDWTKPISELPSLFTSPEYYD
PWHDAVVNNPFSKGFVKFANTYTVRDPQEMFLFQLCEHGESI LYEQTLLALEQTDYCDFEVQ
FEVLHNVIHYLVGGRTYALSSSLHYASYDPFFFIHHSFVDKMWVWQALQKRRKLPYKRAD
CAVNLMTKPMRPFDS DMNQNPFTKMHAVPNTLYDYETLYSYDNLEIGGRNLDQLQAEIDR

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SRSHDRVFAGFLLRGIGTSADVRFWICRNENDCHRGIIIFILGGAKEMPWSFDRNFKFDIT
HVLENAGISPEDVFDAEPPFYIKVEIHAVNKTMI PSSVIPAPTIIYSPGE

Domain g

GRAADSAHSANIAGSGVRKDVTTTLTVSETENLRQALQGVIDDTGPNQYQAIASFHGSPPMC
EMNGRKVACCAHGMA SFPHWHRLYVKQMEDALADHGSHIGIPYWDWTTAFTLPALVTDSE
NNPFHEGRIDHLGVTTSSRS PRDMLFNDPEQGSSEFFYRQVLLALEQTDYCOFEVQFELTHN
AIHSWTGGRSPYGMSTLEFTAYDPLFWLHHSNTDRIWAVWQALQKYRGLPYNEAHCEIQVL
KQPLRPFND D INHNPI TKTNARPIDSFDYERFNYQYDTLSFHGKSIPELNDLLEERKREER
TFAAFLLRGIGCSADVVF DICRPN GDCVFAGTF AVLGGELEMPWSFDRLFRYDITRVMNQL
HLQYDSDFSFRVKLVATNGTELSSDLLKSPTIEHEL

Domain h

GAHRGPVEETEVTROHTDGN AHFHRKEVDSL SLDEAN NLKNALYKLQNDHSLTG YE AISGY
HGYPNLCPEEGDDKI PLLRPRMGIFPYWHRLTIQLERALEHNGALLGVPYWDWNKDLSSL
PAFFSDSSNNNPYFKYHIAGVGHD TVREPTSLIYNQPQIHGYDYLYYLALTTLEENNYWDF
EVQYEILHNAVH SWLGG SOKYS MSTLEYS AFDPVFMILHSGLDRLWIIWQELQKIRRKPIN
FAK CAYHMMEEPLAPFSYPSINQDEFTRANSKPSTVFD SHKFGYHYDNLNVRGHSIQELNT
IINDLRNTDRIYAGFVLSGIGTSASVKIYLRTDDNDEEVGTFTVLGGEREMPWAYERVF KY
DITEVADRLKIKLWGHPLTSGTGDHILTNGIGGKQEP TQILSSSTDLPIMTTMFLLSQXGR
NLHI PPKV VVKGTRIEFHPVDDSVTRPVVDLGSYTALENCVVP PFTYHGFELNHVYSVKP
GDYYVTGPTRDLCQ NADVRIHIHVEDE

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Figure 8

KLH1 cDNA sequence and intron structure

Domain b

GGCCTACCGTACTGGGACTGGACTGAACCCATGACACACATTCCGGGTCTGGCAGGAAACA
AAACTTATGTGGATTCTCATGGTGCATCCACACAAATCCTTTTCATAGTTCAGTGATTGC
ATTTGAAGAAAATGCTCCCCACACCAAAGACAAATAGATCAAAGACTCTTTAAACCCGCT
ACCTTTGGACACCACACAGACCTGTTCAACCAGATTTTGTATGCCTTTGAACAAGAAGATT
ACTGTGACTTTGAAGTCCAATTTGAGATTACCCATAACACGATTTCACGCTTGGACAGGAGG
AAGCGAACATTTCTCAATGTCGTCCCTACATTACACAGCTTTCGATCCTTTGTTTTACTTT
CACCATTCTAACGTTGATCGTCTTTGGGCCGTTTGGCAAGCCTTACAGATGAGACGGCATA
AACCCTACAGGGCCCACTGCGCCATATCTCTGGAACATATGCATCTGAAACCATTTCGCCTT
TTCATCTCCCCTTAACAATAACGAAAAGACTCATGCCAATGCCATGCCAAAACAAGATCTAC
GACTATGAAAATGTCCTCCATTACACATACGAAGATTTAACATTTGGAGGCATCTCTCTGG
AAAACATAGAAAAGATGATCCACGAAAACCAGCAAGAAGACAGAATATATGCCGGTTTTCT
CCTGGCTGGCATACTACTTCAGCAAATGTTGATATCTTCATTAAAACTACCGATTCCGTG
CAACATAAGGCTGGAACATTTGCAGTGCTCGGTGGAAGCAAGGAAATGAAGTGGGGATTTG
ATCGCGTTTTTCAAGTTTGACATCACGCACGTTTTTGAAGATCTCGATCTCACTGCTGATGG
CGATTTCGAAGTTACTGTTGACATCACTGAAGTCGATGGAACATAAATTGCATCCAGTCTT
ATTCCACATGCTTCTGTTCATTTCGTGAGCATGCACGTGGTAAGCTGAATAGAG

Intron b/c

GTTTTGTAATAATTATGTAGAATTCTTTACCTCAGAATAAGATGAGGTCACATGGGTTTTG
CAAACTATTACGTTCGAATTAATATTAATAATACCGGACCCTCCACTGGTACATATTTAT
CTTTATAACGATAATAGCGATGATGATGATGATGATGATGATGATGATGATGATGATgATAATg
ATgATGCCGGTATTGCACGTAATCCAGCCGAcTTAGATGACACCCTAAGGGTGCAGAAAGT
ATAaCAATTAGATTGCGTTtGCATCTGTGTATGCGTGTGCTTTAaCCAAAAGTCAAAATAA
AAGTGCAAACCCTTAGTTTATTTCATTTGATAGAGCCTTTTACGATAAGAACAATGTAATAA
ATTAGAACATAACTGAAACCTCCGAAAGAAGGCCTGTTTGTCAAGAGAGGTATCGACATGA
TTGACTTATAAACCTGTGCTTCTATATTTTGGAACTGTCCACTTTCTTGTTGTGTGTACTG
TAATCACATCGCACTATGGCTGCAAGACGTGTACGAGTACACTATATACTTACCTAATGAC
CAACCACAAGGCTGGCTTTGTTAATATTGTTATTTACAGAAATAAACACAGAATTCCAGC
ATTTGGCTGGTGTATTTAGCAAAACACCGATATGACACTCATGTTTTATTACATTTTTTTTC
AG

Domain c

TTAAATTTGACAAAGTGCCAAGGAGTCGTCTTATTTCGAAAAAATGTAGACCGTTTGAGCCC
CGAGGAGATGAATGAACTTCGTAAAGCCCTAGCCTTACTGAAAGAGGACAAAAGTGCCGGT
GGATTTTCAGCAGCTTGGTGCATTCCATGGGGAGCCAAAATGGTGTCTTAGTCCCGAAGCAT
CTAAAAAATTTGCCTGCTGTGTTTACGGCATGTCTGTGTTCCCTCACTGGCATCGACTGTT
GACGGTTCAGAGTGAAAATGCTTTGAGACGACATGGCTACGATGGAGCTTTGCCGTACTGG
GATTGGACCTCTCCTCTTAATCACCTTCCCGAACTGGCAGATCATGAGAAGTACGTCGACC
CTGAAGATGGGGTAGAGAAGCATAACCCTTGGTTCGATGGTCATATAGATACAGTCGACAA
AACAAACAAGAAGTGTTTCAAGATAAACTCTTCGAACAGCCTGAGTTTGGTCATTATACA
AGCATTGCCAAACAAGTACTGCTAGCGTTGGAACAGGACAATTTCTGTGACTTTGAAATCC
AATATGAGATTGCCATAACTACATCCATGCACTTGTAGGAGGCGCTCAGCCTTATGGTAT
GGCATCGCTTCGCTACACTGCTTTTGATCCACTATTCTACTTGCATCACTCTAATACAGAT
CGTATATGGGCAATATGGCAGGCTTTACAGAAGTACAGAGGAAAACCGTACAACGTTGCTA
ACTGTGCTGTTACATCGATGAGAGAACCTTTGCAACCATTTGGCCTCTCTGCCAATATCAA
CACAGACCATGTAACCAAGGAGCATTTCAGTGCCATTCAACGTTTTTGATTACAAGACCAAT

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TTCAATTATGAATATGACACTTTGGAATTTAACGGTCTCTCAATCTCTCAGTTGAATAAAA
AGCTCGAAGCGATAAAGAGCCAAGACAGGTTCTTTGCAGGCTTCCTGTTATCTGGTTTCAA
GAAATCATCTCTTGTTAAATTCAATATTTGCACCGATAGCAGCAACTGTCACCCCGCTGGA
GAGTTTTACCTTCTGGGTGATGAAAACGAGATGCCATGGGCATACGATAGAGTCTTCAAAT
ATGACATAACCGAAAACTCCACGATCTAAAGCTGCATGCAGAAGACCACTTCTACATTGA
CTATGAAGTATTTGACCTTAAACCAGCAAGCCTGGGAAAAGATTTGTTCAAGCAGCCTTCA
GTCATTCATGAACCAAGAATAG

Intron c/d

GTACTTGTTATATGTTTCGAATATTGCCGATACCTTCAATATATATACTTTATCAAAGTAA
TTGATTAATCTGAAGTAATTTTCCTTTCCAGTAGAGATTCAGTTGATACAACAAGAATTG
CCCTGTTGTATGTCACCTTTATTTTCATCAAACGATTTCGAAGTGAGCTGTCCATGCCACAAT
GGGGTCTCTGTAACCTTTCTCGTATGGGGTATAGATTATATAGACGTGGCAGACCTTACGTA
TAACTAATATTTGTGTAATGTCGTTTCAG

Domain d

GTCACCATGAAGGCGAAGTATATCAAGCTGAAGTAACTTCTGCCAACCGTATTTCGAAAAAA
CATTGAAAATCTGAGCCTTGGTGAACCTCGAAAGTCTGAGAGCTGCCTTCCTGGAAATTGAA
AACGATGGAACCTTACGAATCAATAGCTAAATTCCATGGTAGCCCTGGTTTGTGCCAGTTAA
ATGGTAACCCCATCTCTTGTTGTGTCCATGGCATGCCAACTTTCCCTCACTGGCACAGACT
GTACGTGGTTGTGCTTGAGAATGCCCTCCTGAAAAAAGGATCATCTGTAGCTGTTCCCTAT
TGGGACTGGACAAAACGAATCGAACATTTACCTCACCTGATTTTCAGACGCCACTTACTACA
ATTCCAGGCAACATCACTATGAGACAAACCCATTCCATCATGGCAAAATCACACACGAGAA
TGAAATCACTACTAGGGATCCCAAGGACAGCCTCTTCCATTTCAGACTACTTTTACGAGCAG
GTCCTTTACGCCTTGGAGCAGGATAACTTCTGTGATTTTCGAGATTCAGTTGGAGATATTAC
ACAATGCATTGCATTCTTTACTTGGTGGCAAAGGTAAATATTCCATGTCAAACCTTGATTA
CGCTGCTTTTGATCCTGTGTTCTTCCTTCATCACGCAACGACTGACAGAATCTGGGCAATC
TGGCAAGACCTTCAGAGGTTCCGAAAACGGCCATACCGAGAAGCGAATTGCGCTATCCAAT
TGATGCACACGCCACTCCAGCCGTTTGATAAGAGCGACAACAATGACGAGGCAACGAAAAC
GCATGCCACTCCACATGATGGTTTTGAATATCAAAACAGCTTTGGTTATGCTTACGATAAT
CTGGAAGTGAATCACTACTCGATTCCCTCAGCTTGATCACATGCTGCAAGAAAGAAAAAGGC
ATGACAGATATTTCGCTGGCTTCCTCCTCACAAATATTGGAACatCTGCCGATGGCCATGT
ATTTGTATGTCTCCCAACTGGGGAACACACGAAGGACTGCAGTCATGAGGCTGGTATGTTT
TCCATCTTAGGCGGTCAAACGGAGATGTCCTTTGTATTTGACAGACTTTACAAACTTGACA
TAACTAAAGCCTTGAAAAAGAACGGTGTGCACCTGCAAGGGGATTTTCGATCTGGAAATTGA
GATTACGGCTGTGAATGGATCTCATCTAGACAGTCATGTCATCCACTCTCCCACTATACTG
TTTGAGGCGCGAACAG

Intron d/e

GTAACATTTTTGTCACTGTAACCAACAACCTGCAGTCTATTTTGCAATTACGATAATAACAA
TTTTTGAAATATATCTTTATTAAAGCAAAGGTTTCTAGAGACAAACAGCCGGCTCTAATTA
TTTTTTTCGAACTTACGCTTGAGTAAAGATCTGCAAATGGCAACCCTACCTATACTATTAAA
AATATAATGTTACATTTCGTATCTGAATGTTTAATAAATCACTTCATATTCTGTTGCAG

Domain e

ATTCTGCCCCACACAGATGATGGACACACTGAACCAGTGATGATTCGCAAAGATATCACACA
ATTGGACAAGCGTCAACAACCTGTCACTGGTGAAAGCCCTCGAGTCCATGAAAGCCGACCAT
TCATCTGATGGGTTCCAGGCAATCGCTTCCTTCCATGCTCTTCCTCCTCTTTGTCCATCAC
CAGCTGCTTCAAAGAGGTTTGCGTGCTGCGTCCATGGCATGCCAACCTTCCCGCAATG

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Figure 9

Derived primary structure of KLH1

Domain b

GLPYWDWTEPMTHIPGLAGNKTYVDSHGASHTNPFHSSVIAFEENAPHTKRQIDQRLFKPA
TFGHHTDLFNQILYAFEQEDYCDFEVQFEITHNTIHAWTGGSEHFSMSSLHYTAFDPLFYF
HHSNVDRLWAVWQALQMRHKPYRAHCAISLEHMHLPFAFSSPLNNNEKTHANAMPNKIY
DYENVLHYTYEDLTFFGGISLENIEKMIHENQQEDRIYAGFLLAGIRTSANVDIFIKTTSV
QHKAGTFAVLGGSKEMKWGFDRVFKFDITHVLKDLDLTADGDFEVTVDITEVDGTKLASSL
IPHASVIREHARGKLN

Domain c

VKFDKVPRLIRKNVDRLSPEEMNELRKALALLKEDKSAGGFQQLGAFHGEPKWCPSPEA
SKKFACCVHGMVFPWHRLTLVQSENALRRHGYD GALPYWDWTSPLNHLPELADHEKYVD
PEDGVEKHNPWFDPGHIDTVDKTTTRSVQNKLFEPFGHYTSIAKQVLLALEQDNFCDFEI
QYEIAHNYIHALVGGGAQPYGMASLRYTAFDPLFYLHHSNTDRIWAIWQALQKYRGKPYNVA
NCAVTSMPREPLQPFGLSANINTDHVTKEHSVPFNVFDYKTNFNIEYDTLEFNGLSISQLNK
KLEAIKSQDRFFAGFLLSGFKKSSLVKFNICTDSSNCHPAGEFYLLGDENEMPWAYDRVFK
YDITEKLHDLKLHAEDHFYIDYEVFDLKPASLGKDLFKQPSVIEPRI

Domain d

GHHEGEVYQAEVTSANRIRKNIENLSLGELESLRAAFLEIENDGTYESIAKFHGSPGLCQL
NGNPISCCVHGMPFPHWHRLYVVVENALLKKGSSVAVPYWDWTKRIEHLPHLISDATYY
NSRQHHEYNPFHHGKI THENEITTRDPKDSLFSDFYEQVLYALEQDNFCDFEIQLEIL
HNALHSLGKGKYSMSNLDYAAFDPVFFLHHATTDRIWAIWQDLQRFKRPYREANCAIQ
LMHTPLQPFDKSDNDEATKTHATPHDGFYQNSFGYAYDNLELNHYSIPQLDHMLQERKR
HDRVFAGFLLHNIGTSADGHVFCVCLPTGEHTKDCSHEAGMFSILGGQTEMSFVFDRLYKLD
ITKALKKNGVHLQGDFFLEIEITAVNGSHLDSHVIHSPTILFEAG

Domain e

DSAHTDDGHTEPVMIRKDI TQLDKRQQLSLVKALESMKADHSSDGFQAIASFHALPPLCPS
PAASKRFACCVHGMPFPQWHRLYTVQFQDSL RKHGAVVGLPYWDWTLPR

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Figure 10

KLH2 cDNA sequence and intron structure

Domain b

GGCCTGCCCTACTGGGATTGGACCATGCCAATGAGTCATTTGCCAGAACTGGCTACAAGTG
 AGACCTACCTCGATCCAGTTACTGGGGAACTAAAAACAACCCTTTCCATCACGCCCCAAGT
 GGCGTTTGAAAATGGTGTAAACAAGCAGGAATCCTGATGCCAACTTTTTATGAAACCAACT
 TACGGAGACCACACTTACCTCTTCGACAGCATGATCTACGCATTTGAGCAGGAAGACTTCT
 GCGACTTTGAAGTCCAATATGAGCTCACGCATAATGCAATACATGCATGGGTTGGAGGCAG
 TGAAAAGTATTCAATGTCTTCTCTTCACTacacTGCTTTTGATCCTATATTTTACCTCCAT
 CACTCAAATGTTGATCGTCTCTGGGCCATTTGGCAAGCTCTTCAAATCAGGAGAGGCAAGT
 CTTACAAGGCCCACTGCGCCTCGTCTCAAGAAAGAGAACCATTAAAGCCTTTTGCATTTCAG
 TTCCCCACTGAACAACAACGAGAAAACGTACCACAACCTCTGTCCCCACTAACGTTTATGAC
 TATGTGGGAGTTTTGCACTATCGATATGATGACCTTCAGTTTGGCGGTATGACCATGTGAG
 AACTTGAGGAATATATTACACAAGCAGACACAACATGATAGAACCTTTGCAGGATTCTTCCT
 TTCATATATTGGAACATCAGCAAGCGTAGATATCTTCATCAATCGAGAAGGTCATGATAAA
 TACAAAGTGGGAAGTTTTGTAGTACTTGGTGGATCCAAAGAAATGAAATGGGGCTTTGATA
 GAATGTACAAGTATGAGATCACTGAGGCTCTGAAGACGCTGAATGTTGCAGTGGATGATGG
 GTTCAGCATTACTGTTGAGATCACCGATGTTGATGGATCTCCCCCATCTGCAGATCTCATT
 CCACCTCCTGCTATAATCtTTGaACGTGGTCaTG

Intron 2b/c

AGGTATTTAAAAAAGTAATAAAACCATATTTTCGAATGCGCTTTATGAAATATCGTGTGAC
 TGGTTCTTTAGTTTACATGGAGTGTAACAACATGCTCCATCAGTTGACATATACTGCTCAC
 ACAAGTAAGGGATATTTGATAATGATAACAAATATAATCAAAGCGGTTATACTATCAAGA
 CTTATTACATAATTACAGGTGAAGGGAGGTGTGATCGTGTTCACTGATCAGGTTGAGGCC
 AGAGAAAGTCCAGTTTGAGTCTTGCAAGAGATGATGTTTAGGCATGGGGTCGAATCACCBA
 AATCACATGACTTCAATAACGGGTTGGACCACCTCGAGCGACgATGCAAGCAGTAGAGCGT
 CTACGCATGCTCCTGATAAGGCGACCAATCTGTTCCCTGGGGAATCAGtCGCCACTCCTCTT
 GTAGTGCCACGCTCATTCTGCTACGGTCCTGGGTACCTGCTATCGGgTCTTGATCCGTAT
 CCCAAGGATGTCCACACATGTTCAAgGTGAGAGGTGCGGGAACATCGCTGGCCACGGTtA
 GGTCTGAATTTGATGCCGTTGAAAGTGAGCTCTGACAACcTGAGCATGGtGAGCTCTGACG
 TTGTGCTCCTGAAAGATGAATcCAGCTcCaTGaCAGCGAGCAAaGGGCAGGACGTGTTGGT
 CAATGCAGTTGTCTCTGCAGTACACACCTGTCACTCGCCACTCACAAGCGTGTAGATCTGT
 ACGACCAGTCATGGAGATCCCAGCCCACATCATAACGGACCCCTATCCATACCGATCATGA
 GCCACCATAGCAGCGTCTTGATGACGTTCTCCCTGTGCGCTCGACATCCTcACACGGCCAA
 AAGGAACGTGGACTCGTCACTGAACATGACATTAGCCAACCTGGCACTTGTCCACCGCTGA
 TGTGGCGAGACCATTCCAGTCGAGCTCTTCGGTGTCTGGCTTTTCATCGATAACACGACGT
 AAGGTCTGCGGGCGTGAAGACGGCTCTATGCAGGCGATTTCGGATTGTCTGGGTGCTAAC
 TCTGATCCCAGGTGCCTGCTGAAGTTGATGCTGGATCTGTGTGGCATTGAGATGGCGATT
 CTTAGGACTGTGGAGATGATGAATCGATCTTGACTTATGGTGGTGACATTAGGACGTCGGG
 TTCGTGTCCTATCCTGCACCTCTTCCAGTTGTTTCGGTGACGCTCTGGTACCCGGCTGATTAC
 TGACTGAGAATATCCATCTGCCGTGCGACATGAGCCTGTGTTGGCCCAGCCTGAAGCATTG
 CAATCGCCAGAGACGCTCTTCAAAAGTCATTCGACGCATGGtTTTCTGTTACAAATGACA
 GCGTAAAACAGtTTTTGGtGCTTTTATGCTTCCCAAGAGCATGAAAAACACGTTCTATgGG
 TCGtGCACACCTTACATGACAAGtGtGAAAAGtGACTTGcACCCCCCTTGTGtGTTTCGGATG
 CACACTCTGTTTACGTACTGATGCGATTTGGCGTCTAAACATGTTTTGGCGTCTAAACATG
 TTTTCCTGCATGATTCATATACTATTTTGTGATATTCCTGGCATCAAACCAAACCTACAGTG
 AAATATATTTCAATATCCCCTACTTTGTGTGAGTAGTATAGATCACTGCAGACAACATATA

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GACAAtGCAGtTaCaCCGTCAACAATCCCAGTCATTAATTATGATGaCaCTTCCACACATA
GIGTCAGTGATTGTAATTCAaCTGTACACACTTTTCCCGTGAACATTTCAGGATCTATATGA
CTAAATATATAACATTAGTATACGTGCAGTTTTGTATCGCTACGACATTGTTGTAACCTCTT
TGTTTAATCATTTaACAG

Domain c

CTGATGCCAAAGaCTTTGgCCATAGCAGAAAAATCAGgAAAGcCGTTGATTcTcTGACAGT
CgAAGAACAAAcTTCGTTGAGgCGAGcTATGgCAGATcTACAGGACGACAAAACATCAGGG
GGTTTCCAGCAGATTGCAGCATTCACGGAGAACC AAAATGGTGTCCAAGCCCCgAAGCGG
AGAAAAAATTTGCATGCTGTGTTTCATGGAATGGCTGTTTTCCCTCACTGGCACAGATTGCT
GACAGTTCAAGGAGAAAAATGCTCTGAGGAAACATGGCTTTACTGGTGGACTGCCCTACTGG
GACTGGACTCGATCAATGAGCGCCCTTCCACATTTTGTGCTGATCCTACTTACAATGATG
CTATTTCCAGCCAGGAAGAAGATAACCCATGGCATCATGGTCACATAGACTCTGTTGGGCA
TGATACTACAAGAGATGTGCGTGATGATCTTTATCAATCTCCTGGTTTCGGTCACTACACA
GATATTGCACAACAAGTCCTTCTGGCCTTTGAGCAGGACAGTTTCTGTGATTTTGAGGTAC
AATTTGAAATTGCCATAATTTATACATGCACTGATTGGTGGAACGAACCATACAGTAT
GTCATCTTTGAGGTATACATACGATCCAATCTTCTTCTTGCAACCACTCCAGTACAGAC
CGACTTTGGGCCATCTGGCAAGCAATCACTAGTGCGGCCGCCCTGCAGGTGACCATAAAGG
AGAGCTCCCAACGCgtTGGAtGCAATCT

Domain g

ATGGCTGTGTTTCCGCACTGGCACAGACTGTTTGTGAAACAGATGGAGGACGCACTTGCTG
CTCATGGAGCTCATATTGGCATACCATACTGGGATTGGACAAGTGCGTTTAGTCATCTGCC
CGCCCTAGTGACTGACCACGAGAACAATCCCTTCCACCAC

Intron g(2)

GTATGTGTCAAATCGTTTTAGGAACTGCCTTATCCATTTTACAATTACGAGTACAAAATGA
AAACGGAAACTGTGTGACCTCGAAAAGTGCAATCTTTAAAGGATGCAATGTACACAATAAA
ATGCTCCGATCAAAGCGATGGCTAGAAATCATTTTCCCTCTAATTCCCTTTCACACAGCT
CGGTTTCGTTTTAAGTAGGAACAAGTCTCTGCAAAAACATCACAAATAAAGAGAACACAGAA
AAAACCTCATTCTCGTTTCTGTATTCCGAAAATGAAATTTACAATTTCTTTTCAATTTATAG

Domain g

GGCCATATTGGTCATCTGAATGTGGATACATCTCGATCTCCAAGAGACATGCTGTTTAATG
ATCCTGAACAAGGCTCAGAATCATTCTTCTACAGACAGGTCTCTTGACTCTAGAACAGAC
AGACTTCTGCCAATTTGAAGTTCAGTTTGAACCTTACACACAATGCCATCCACTCTTGGACT
GGAGGACATACTCCATATGGAATGTCATCACTGGAATATACAGCATATGATCCACTCTTTT
ATCTCCACCATTCCAACACTGATCGTATCTGGGCCATCTGGCAGGCACTCCAGAAATATAG
AGGTCTTCCATACAACGCAGCTCACTGCGATATccaagttctgaaacaacctCTTAAACCA
TTCAGCGAGTCCAGGAATCCAAACCCAGTCACCAGAGCCAATTCTAGGGCCGTTGATTCAT
TTGATTATGAGAAATTCAATTATCAATATGACACACTTACCTTCCACGGACTTCTATCCC
AGAACTTGATGCCATGCTTCAAGAGAGAAAGAAGGAAGAGAGAACATTTGCAGCCTTCCTG
TTGCACGGATTTGGCGCCAGTGCTGATGTTTCGTTTGATGTCTGCACACCTGATGGTCATT
GTGCCTTTGCTGGAACCTTCGCGGTACTTGGTGGGGAGCTTGAGATGCCCTGGTCCTTTGA
AAGATTGTTCCGTTACGATATCACAAAGGTTCTCAAGCAGATGAATCTTCACTATGATTCT
GAGTTCCACTTTGAGTTGAAGATTGTTGGCACAGATGGAACAGAAGTCCCATCGGATCGTA
TCAAGAGCCCTACCATTGAACACCATGGAGGAG

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Intron g/h

GTATGTTTTGAGATCCACATAATCTTCTACCCTGTCTCATTTCTAATGCTCTTCAATACAC
AATTTATATAGCCTTTGAGCTTCAGATGTATTACGGACAGGCATTACAGTATACATGTAAT
ATGGTTTTCTGCTATTTGCAAAAATTGTGTCCTATCTCTGTTTCAGATCATCATGGCGGTGA
CACCTAG

Domain h

GTCACGATCACAGTGAACGTCACGATGGATTTTTTCAGGAAGGAAGTCGGTTCCCTGTCCCT
GGATGAAGCCAATGACCTTAAAAATGCACTGTACAAGCTGCAGAATGATCAGGGTCCCAAT
GGATATGAATCAATAGCCGGTTACCATGGCTATCCATTCTCTGCCCTGAACATGGTGAAG
ACCAGTACGCATGCTGTGTCCACGGAATGCCTGTATTTCCACATTGGCACAGACTTCATAC
AATCCAGTTTGAGAGAGCTCTCAAAGAACATGGTTCTCATTTGGGTCTGCCATACTGGGAC
TGGAC

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Figure 11

Derived primary structure of KLH2

Domain b

GLPYWDWTMPMSHLP ELATSETYLD PVTGETKNNPFHHAQVAFENGVT SRNPDAKLFMKPT
YGDHTYLFDSMIYAFEQEDFCDFEVQYELTHNAIHAWVGGSEKYSMSLHYTAFDPIFY LH
HSNVDRLWAIWQALQIRRGKSYKAHCASSQEREPLKPFASFSSPLNNNEKTYHNSVPTNVYD
YVGVLHYRYDDLQFGGMTMSELEEYIHKQTQHDRTFAGFFLSYIGTSASVDIFINREGHDK
YKVGSEFVVLGGSKEMKWGFDRMYKYEITEALKTLNVAVDDGFSITVEITDVDGSPPSADLI
PPPAIIFERGH A

Domain c

DAKDFGHSRKIRKA VDSLTVEEQTS LR RAMADLQDDKTSGGFQQIAAFHGE PKWCPSPEAE
KKFACCVHGM AVFPHWHRLLT VQGENALRKHGFTGGLPYWDWTRSMSALPHFVADPTYNDA
ISSQEEDNPWHHGHIDSVGHDTTRDVRDDLQSPGFGHYTDIAQQVLLAFEQDSFCDFEVQ
FEIAHNFIHALIGGNEPYSMSLRYTTYDPIFFLHHSSTDRLWAIWQALQKYRGKPYNTAN
CAIASMRKPLQPFGLDSVINPDDETREHSVPFRVFDYKNNFDY EYESLAFNGLSIAQLDRE
LQRRKSHDRVFAGFLLHEIGQSAKHNVSDCDHYAGEFYILGDEAEMPWRYDRVYKYEITQQ
LHDLDLHVGDNFFLKYEA FDLNGGSLGGSIFSQPSVIFEPAAGMF

Domain d

GSHQADEYREAVTSASHIRKNIRD LSEGEIESIRSAFLQIQKEGIYENIAKFHGKPGLC EH
DGHPVACCVHGMPTFPHWHRLYLQVENALLERGS AVAVPYWDWTLPR

Domain g

MAVFPWHRL FVKQMEDALAAHGAHIGIPYWDWTS AFSHLPALVTDHENNPFHHGHIGHLN
VDTSRSPRDM LFNDPEQGSSEFFYRQVLLTLEQTD FCQFEVQFELTHNAIHSWTGGHTPYG
MSSLEYTAYDPLFY LHHSNTDRIWAIWQALQKYRGLPYNAAHCDIQVLKQPLKPFSESRNP
NPVTRANSRAVDSFDYEKFNYQYDTLT FHGLSIPELDAMLQERKKEERTFAAFL LHGFGAS
ADV SFDVCTPDGHCAFAGTFAVLGGELEMPWSFERLFRYDITKVLKQMN LHVDSEFHFELK
IVGTDGTELP SDRIKSPTIEHHGG

Domain h

GHDHSE RHDGFFRKEVGSLSLDEANDLKNALYKLQNDQGPNGYESIAGYHGY PFLCPEHGE
DQYACCVHGM PVFPHWHRLHTIQFERALKEHGSHLGLPYWDW